

Re-RUN



RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/596,141C

DATE: 03/07/2003 TIME: 10:46:51

Input Set : N:\vernette\US09596141C.raw
Output Set: N:\CRF4\03072003\I596141C.raw

```
1 <110> APPLICANT: Lawn, Richard M.
         Wade, David
 3
         Oram, John F.
         Garvin, Michael
 4
 5 <120> TITLE OF INVENTION: Compositions and Methods for Increasing Cholesterol
         Efflux and Raising HDL using ATP Binding Cassette
 6
 7
         Transporter Protein ABC1
 8 <130> FILE REFERENCE: 99,395-B
9 <140> CURRENT APPLICATION NUMBER: US/09/596,141C
10 <141> CURRENT FILING DATE: 2000-06-16
11 <150> PRIOR APPLICATION NUMBER: US 60/140,264
12 <151> PRIOR FILING DATE: 1999-06-18
13 <150> PRIOR APPLICATION NUMBER: US 60/153,872
14 <151> PRIOR FILING DATE: 1999-09-14
                                                          ENTERED
15 <150> PRIOR APPLICATION NUMBER: US 60/166,573
16 <151> PRIOR FILING DATE: 1999-11-19
17 <160> NUMBER OF SEQ ID NOS: 62
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20 <211> LENGTH: 10442
21 <212> TYPE: DNA
22 <213> ORGANISM: Homo sapiens
23 <220> FEATURE:
24 <221> NAME/KEY: unsure
25 <222> LOCATION: (1)..(10442)
26 <223> OTHER INFORMATION: All n's are unknown.
27 <400> SEQUENCE: 1
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actctacatc tecetteecq ageaaggage tggetgaage cacaaaaaca ttgetgeata 1080 45 46 qtcttqqqac tctqqcccaq qaqctqttca qcatqaqaaq ctqqaqtqac atgcgacagg 1140 aggtgatgtt tctgaccaat gtgaacagct ccagctcctc cacccaaatc taccaggctg 1200 47 tqtctcqtat tqtctqcqqq catcccqaqq qaqqqqqqct qaaqatcaaq tctctcaact 1260 48 49 ggtatgagga caacaactac aaagccctct ttggaggcaa tggcactgag gaagatgctg 1320 aaaccttcta tgacaactct acaactcctt actgcaatga tttgatgaag aatttggagt 1380 50 ctaqtcctct ttcccqcatt atctqqaaaq ctctqaaqcc gctqctcgtt gggaagatcc 1440 51 tgtatacacc tgacactcca gccacaaggc aggtcatggc tgaggtgaac aagaccttcc 1500 52 53 aggaactggc tgtgttccat gatctggaag gcatgtggga ggaactcagc cccaagatct 1560 qqaccttcat qqaqaacagc caaqaaatgg accttgtccg gatgctgttg gacagcaggg 1620 54 55 acaatgacca cttttgggaa cagcagttgg atggcttaga ttggacagcc caagacatcg 1680 56 ggagagaagc tttcaacgag actaaccagg caatccggac catatctcgc ttcatggagt 1800 57 gtgtcaacct gaacaagcta gaacccatag caacagaagt ctggctcatc aacaagtcca 1860 58 tqqaqctqct qqatqaqaqq aaqttctqqq ctqqtattqt gttcactgga attactccag 1920 59 qcaqcattqa qctqccccat catqtcaaqt acaaqatccq aatgqacatt gacaatgtgg 1980 60 agaggacaaa taaaatcaag gatgggtact gggaccctgg tcctcgagct gacccctttg 2040 61 aggacatgcg gtacgtctgg gggggcttcg cctacttgca ggatgtggtg gagcaggcaa 2100 62 tcatcagggt gctgacgggc accgagaaga aaactggtgt ctatatgcaa cagatgccct 2160 63 atccctgtta cgttgatgac atctttctgc gggtgatgag ccggtcaatg cccctcttca 2220 64 65 tgacgctggc ctggatttac tcagtggctg tgatcatcaa gggcatcgtg tatgagaagg 2280 aggcacggct gaaagagacc atgcqqatca tgggcctgga caacagcata ctctggttta 2340 66 67 qctqqttcat tagtagcctc attcctcttc ttgtgagcgc tggcctgcta gtggtcatcc 2400 tgaagttagg aaacctgctg ccctacagtg atcccagcgt ggtgtttgtc ttcctgtccg 2460 68 tgtttgctgt ggtgacaatc ctgcagtgct tcctgattag cacactcttc tccagagcca 2520 69 70 acctggcagc agcctgtggg ggcatcatct acttcacgct gtacctgccc tacgtcctgt 2580 qtqtqqcatq qcaqqactac gtqgqcttca cactcaagat cttcgctagc ctgctgtctc 2640 71 ctgtggcttt tgggtttggc tgtgagtact ttgccctttt tgaggagcag ggcattggag 2700 72 tgcagtggga caacctgttt gagagtcctg tggaggaaga tgqcttcaat ctcaccactt 2760 73 cgatctccat gatgctgttt gacaccttcc tctatggggt gatgacctgg tacattgagg 2820 74 75 ctgtctttcc aggccagtac ggaattccca ggccctggta ttttccttgc accaagtcct 2880 76 actggtttgg cgaggaaagt gatgagaaga gccaccctgg ttccaaccag aagagaatgt 2940 cagaaatctg catggaggag gaacccaccc acttgaagct gggcgtgtcc attcagaacc 3000 77 tggtaaaagt ctaccgagat gggatgaagg tggctgtcga tggcctggca ctgaattttt 3060 78 79 atgagggcca gatcacctcc ttcctgggcc acaatggagc ggggaagacg accaccatgt 3120 caateetgae egggttgtte eeceegaeet egggeaeege etacateetg ggaaaagaea 3180 80 ttcgctctga gatgagcacc atccggcaga acctgggggt ctgtccccag cataacgtgc 3240 81 tqtttqacat qctqactqtc qaagaacaca tctqgttcta tgcccgcttg aaagggctct 3300 82 ctqaqaaqca cqtqaaqqcq qaqatqqaqc agatgqccct ggatgttggt ttgccatcaa 3360 83 gcaagctgaa aagcaaaaca agccagctgt caggtggaat gcagagaaag ctatctgtgg 3420 84 cettggcctt tgtcggggga tctaaggttg tcattctgga tgaacccaca gctggtgtgg 3480 85 accettacte eegeaggga atatgggage tgetgetgaa atacegacaa ggeegeacea 3540 86 87 ttattctctc tacacaccac atggatgaag cggacgtcct gggggacagg attgccatca 3600 tctcccatgg gaagctgtgc tgtgtgggct cctccctgtt tctgaagaac cagctgggaa 3660 88 89 caqqctacta cctgaccttg gtcaagaaag atgtggaatc ctccctcagt tcctgcagaa 3720 acagtagtag cactgtgtca tacctgaaaa aggaggacag tgtttctcag agcagttctq 3780 90 atgctggcct gggcagcgac catgagagtg acacgctgac catcgatgtc tctgctatct 3840 91 ccaacctcat caggaagcat gtgtctgaag cccggctggt ggaagacata gggcatgagc 3900 92 tgacctatgt gctgccatat gaagctgcta aggagggagc ctttgtggaa ctctttcatg 3960 93

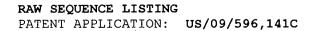
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Input Set : N:\vernette\US09596141C.raw
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169			tgtcttgtgg				
170			aattgcaaca				
171			tcaacaggga				
172	-	-	tttagtacct		_		
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174			atatggagca				
175			gtaaaaatac				
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182			taaaatcgta				
183			ctctgatttc				
184			acacgacatt				
185			atacatcttc				
186			taaacctaat				
187	caatcaagca	adatttctgt	atattccctg	Lygaatgtac	clargigagt	cccayadatt	3000
188			atttctgctt				
189			aaatacagaa				
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Output Set: N:\CRF4\03072003\I596141C.raw

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	209	\400 /				Trp	Pro	Gln	Leu	Arg	Leu	Leu	Leu	Trp	Lys	Asn	Leu	Thr	
	210		_ 1	_	_	_	5		_		_	10	_	~1	1	~ 7	15	_	
	211 212		Phe	Arg	Arg	Arg 20	GIn	Thr	Cys	GIn	Leu 25	Leu	Leu	GLU	vaı	A1a 30	Trp	Pro	
	213		Leu	Phe	Ile		Leu	Ile	Leu	Ile		Val	Arg	Leu	Ser		Pro	Pro	
	214			0.2	35					40					45		_		,
	215 216		Tyr	Glu 50	Gln	His	Glu	Cys	His 55	Phe	Pro	Asn	Lys	Ala 60	Met	Pro	Ser	Ala	
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	218		65				•	70		-			75					80	
	219 220		Cys	Phe	Arg	Tyr	Pro 85	Thr	Pro	Gly	Glu	Ala 90	Pro	Gly	Val	Val	Gly 95	Asn	
	221		Phe	Asn	Lys	Ser		Val	Ala	Arg	Leu	Phe	Ser	Asp	Ala	Arg	Arg	Leu	
	222					100					105					110			
	223 224		Leu	Leu	Tyr 115	Ser	Gln	Lys	Asp	Thr 120	Ser	Met	Lys	Asp	Met 125	Arg	Lys	Val	
	225		Leu	Ara		Leu	Gln	Gln	Ile		Lvs	Ser	Ser	Ser		Leu	Lys	Leu	
	226			130					135	_	_			140			-		
	227			Asp	Phe	Leu	Val	_	Asn	Glu	Thr	Phe		Gly	Phe	Leu	Tyr		
	228		145	-		.	D	150	0	m1.	77 - 7	70	155		T	70	70.7 -	160	
	229 230		Asn	Leu	Ser	Leu	165	Lys	Ser	Thr	Val	170	Lys	Met	Leu	Arg	Ala 175	Asp	
	231		Val	Ile	Leu	His		Val	Phe	Leu	Gln		Tvr	Gln	Leu	His	Leu	Thr	
	232					180	-1-				185	1	- 1			190			
	233		Ser	Leu	Cys	Asn	Gly	Ser	Lys	Ser	Glu	Glu	Met	Ile	Gln	Leu	Gly	Asp	
	234				195					200					205				
	235		Gln		Val	Ser	Glu	Leu	_	Gly	Leu	Pro	Lys		Lys	Leu	Ala	Ala	
	236 237		ΔΙς	210	Δνα	Val	Lou	Δrα	215 Ser	Δερ	Me+	Δερ	Tle	220	Luc	Pro	Ile	I.eu	
	238		225	GIU	ALG	val	neu	230	Ser	noii	rie (rsh	235	ьeu	пуз	110	116	240	
	239			Thr	Leu	Asn	Ser		Ser	Pro	Phe	Pro		Lvs	Glu	Leu	Ala		
	240		- 9				245					250		1-			255		
	0.44			 1	-	era i	-	-		~	-	~ 1		-	TO 7	~ 1	~ 1	T	

Ala Thr Lys Thr Leu Leu His Ser Leu Gly Thr Leu Ala Gln Glu Leu

241

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/09/596,141C

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 9881,9884,9999,10046
Seq#:4; N Pos. 8,19,69,80,104,142,175,191,220
Seq#:6; N Pos. 2805,2808,2923,2970
Seq#:7; N Pos. 9913,9916,10031,10078
Seq#:9; N Pos. 9913,9916,10031,10078
Seq#:59; N Pos. 3,10

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:59,60,61,62

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Output Set: N:\CRF4\03072003\I596141C.rav

L:9 M:270 C: Current Application Number differs, Wrong Format

L:192 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:9840

M:341 Repeated in SeqNo=1

L:537 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0

M:341 Repeated in SeqNo=4

L:646 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:2760

M:341 Repeated in SeqNo=6

L:832 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:9900

M:341 Repeated in SeqNo=7

L:1307 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:9900

M:341 Repeated in SeqNo=9

L:2107 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59 after pos.:0